SEQUENCE LISTING

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<110> Dempcy, Robert O.
     Gall, Alexander A.
     Lokhov, Sergey G.
     Afonina, Irina A.
      Singer, Michael J.
      Kutyavin, Igor V.
      Vermeulen, Nicolaas M.J.
      Epoch Biosciences, Inc.
<120> T-m Leveling Methods
<130> 17682A-003630US
<140> US 10/032,307
<141> 2001-12-21
<150> US 09/054,830
<151> 1998-04-03
<150> US 09/054,832
<151> 1998-04-03
<150> US 09/431,385
<151> 1999-11-01
<150> US 60/186,046
<151> 2000-03-01
<150> US 09/640,953
<151> 2000-08-16
<150> US 09/724,959
<151> 2000-11-28
<150> US 09/796,988
<151> 2001-02-28
<160> 90
<170> PatentIn Ver. 2.1
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<223> Description of Artificial Sequence:minor groove
      binder (MGB) - modified FAM probe
<220>
<221> modified_base
<222> (1)
<223> n = c modified by FAM
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<220>
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<222> (18)
<223> n = t modified by a quencher (Q) and minor groove
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<400> 1
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nttttgacct aacaaatn
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      binder (MGB)-modified FAM probe complement
<400> 2
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atgttaattt gttaggtcaa aagaaaaatc tttaga
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<223> Description of Artificial
      Sequence: 4-amino-3-(prop-1-ynyl)pyrazolo[3,4-d]pyrimidine
      (PPPA) analog of adenosine and pyrazolo[3,4-d]pyrimidine
      analog of guanosine (PPG) containing minor groove binder
      (MGB) -modified FAM probe
                                             Ì,
<400> 3
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tacaattaaa caatccagtt ttctttttag aaatct
<210> 4
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<223> Description of Artificial
      Sequence: 4-amino-3-(prop-1-ynyl)pyrazolo[3,4-d]pyrimidine
      (PPPA) analog of adenosine and pyrazolo[3,4-d]pyrimidine
      guanosine (PPG) containing minor groove binder
      (MGB) -modified FAM probe complement
<220>
<221> modified base
<222> (1)
<223> n = pyrazolo[3,4-d]pyrimidine analog of guanosine
      modified by FAM
<220>
<221> modified base
<222> (5)..(6)
<223> n = pyrazolo[3,4-d]pyrimidine analog of guanosine
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<220>
<221> modified base
<222> (9)..(11)
<223> n =
       4-amino-3-(prop-1-ynyl)pyrazolo[3,4-d]pyrimidine
       analog of adenosine
<220>
<221> modified_base
<222> (15)
<223> n = a modified by a quencher (Q) and minor groove
      binder (MGB)
<400> 4
nttanntcnn nagan
                                                                    15
<210> 5
<211> 21
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: TM-Invader
      probe substituted with six
      pyrazolo[3,4-d]pyrimidine analogs of guanosine
<220>
<221> modified base
<222> (2)..(7)
<223> n = pyrazolo[3,4-d]pyrimidine analog of guanosine
<400> 5
tnnnnncct tggcggctac g
                                                                    21
<210> 6
<211> 21
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<223> Description of Artificial Sequence: TM-Invader
      probe substituted with one
      pyrazolo[3,4-d]pyrimidine analog of guanosine
<220>
<221> modified_base
<222> (5)
<223> n = pyrazolo[3,4-d]pyrimidine analog of guanosine
<400> 6
tgggnggcct tggcggctac g
                                                                    21
<210> 7
<211> 21
<212> DNA
<213> Artificial Sequence
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<220>
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      probe
<400> 7
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<211> 10
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<223> Description of Artificial Sequence:complementary
      target 1
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tcggcggcgt
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      target 2
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acagcggcgt
                                                                    10
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<223> Description of Artificial Sequence:complementary
      target 3
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acagcgacgt
                                                                    10
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<223> Description of Artificial Sequence:complementary
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tcagtgacga
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<223> Description of Artificial Sequence:complementary
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tcagtgacaa
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<223> Description of Artificial Sequence:complementary
      target 6
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                                                                     10
tcaatgacag
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<223> Description of Artificial Sequence:complementary
      target 7
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                                                                     10
acaatgataa
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<223> Description of Artificial Sequence:complementary
      target 8
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ccaataataa
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<223> Description of Artificial Sequence:complementary
      target 9
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gtaataataa
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<210> 17
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<400> 17
aaagttatgt ctacttacag aaa
                                                                    23
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aaagctatgt ctacttacag aaa
                                                                    23
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<223> Description of Artificial Sequence:probe sequence
<400> 19
                                                                   23
aaagtcatgt ctacttacag aaa
<210> 20
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aaagttgtgt ctacttacag aaa
                                                                   23
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<213> Artificial Sequence

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	5	
•	400 01	
	<400> 21	_
	aaagttacgt ctacttacag aaa 2	23
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	<211> 23	
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	•	
	<220>	
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	6	
	· ·	
	400 20	
	<400> 22	
	aaagttatat ctacttacag aaa 2	23
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	7	
	·	
	<400> 23	
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	aaagttatgc ctacttacag aaa 2	. 3
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	8	
	<400> 24	
		:3
	addy categoria and and and and and and and and and an	
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aaagttatgt ctgcttacag aaa
                                                                    23
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aaagttatgt ctatttacag aaa
                                                                    23
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aaagttatgt ctacctacag aaa
                                                                    23
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      13
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aaagttatgt ctactcacag aaa
                                                                    23
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<211> 23
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<223> Description of Artificial Sequence:probe sequence
      14
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<400> 30
aaagttatgt ctacttgcag aaa
                                                                    23
<210> 31
<211> 15
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<213> Artificial Sequence
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<223> Description of Artificial Sequence:target sequence
<400> 31
gtaagtagac ataac
                                                                    15
<210> 32
<211> 15
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<223> Description of Artificial Sequence:target sequence
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<221> modified_base
<222> (1)..(15)
<223> n = 4-amino-3-(prop-1-ynyl)pyrazolo[3,4-d]pyrimidine
      analog of adenosine
<400> 32
gtnngtngnc ntnnc
                                                                    15
<210> 33
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<212> DNA
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<220>
<221> modified_base
<222> (15)
<223> n = c modified by minor groove binder (MGB)
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gtaagtagac ataan
                                                                    15
<210> 34
<211> 15
<212> DNA
<213> Artificial Sequence
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<220>
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<220>
<221> modified base
<222> (1)..(14)
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      analog of adenosine
<220>
<221> modified_base
<222> (15)
<223> n = c modified by minor groove binder (MGB)
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gtnngtngnc ntnnn
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<223> Description of Artificial Sequence:duplex
      complement match
<400> 35
                                                                    10
agctgtgact
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<212> DNA
                                            1
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<223> Description of Artificial Sequence:duplex
      complement 1
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agctgtgact
<210> 37
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<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence:duplex
      complement 2
<400> 37
                                                                    10
agcggtgact
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<210> 38
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<212> DNA
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<220>
<223> Description of Artificial Sequence:duplex
      complement 3
<400> 38
agccgtgact
                                                                     10
<210> 39
<211> 10
<212> DNA
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<223> Description of Artificial Sequence:duplex
      complement 4
<400> 39
agcagagact
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<210> 40
<211> 10
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence:duplex
      complement 5
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agcagggact
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<210> 41
<211> 10
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence:duplex
      complement 6
<400> 41
agcagcgact
                                                                     10
<210> 42
<211> 10
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:duplex
      complement 7
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agcaatgact
                                                                     10
<210> 43
<211> 10
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:duplex
      complement 8
<400> 43
agcattgact
                                                                     10
<210> 44
<211> 10
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:duplex
      complement 9
<400> 44
agcactgact
                                                                    10
<210> 45
<211> 10
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:duplex
      complement match
<400> 45
aataataacc
                                                                    10
<210> 46
<211> 10
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence:duplex
      complement 10
<400> 46
aattataacc
                                                                    10
<210> 47
<211> 10
<212> DNA
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<213> Artificial Sequence

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<223> Description of Artificial Sequence:duplex
      complement 11
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aatgataacc
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<210> 48
<211> 10
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence:duplex
      complement 12
<400> 48
aatcataacc
                                                                    10
<210> 49
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<212> DNA
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<223> Description of Artificial Sequence:duplex
      complement 13
<400> 49
aataaaaacc
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<210> 50
<211> 10
<212> DNA
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      complement 14
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aataagaacc
                                                                    10
<210> 51
<211> 10
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<213> Artificial Sequence
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<223> Description of Artificial Sequence:duplex
      complement 15
<400> 51
aataacaacc
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<220>
 <223> Description of Artificial Sequence:primer
       extension template
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aaccactctg tccta
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ctgtaagtag atataac
                                                                    17
<210> 54
<211> 14
<212> DNA
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ggcaagatat atag
                                                                    14
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<211> 14
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gtgacgcaga ttcc
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gtaagtagac ataac
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cagggagctt tgga
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cactcgtgaa gctg
<210> 59
<211> 15
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<400> 59
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gtaagtaggc ataac
<210> 60
<211> 14
<212> DNA
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<223> Description of Artificial Sequence:probe sequence
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                                                                    14
ccggatgtag gatc
<210> 61
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<212> DNA
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<400> 61
                                                                    14
gattacctgg attt
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<210> 62
 <211> 14
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<213> Artificial Sequence
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<223> Description of Artificial Sequence:probe sequence
<400> 62
ccgtcaatgg tcac
                                                                    14
<210> 63
<211> 12
<212> DNA
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<400> 63
cagcacgtag cc
                                                                    12
<210> 64
<211> 14
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<223> Description of Artificial Sequence:probe sequence
<400> 64
cggctacgtg ctgg
                                                                    14
<210> 65
<211> 14
<212> DNA
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<400> 65
cggctacatg ctgg
                                                                    14
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ctaaatctgc cg
                                                                    12
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 <223> Description of Artificial Sequence:probe sequence
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 tctggatgat gggca
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gttcatgggt gtaat
                                                                     15
<210> 69
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<212> DNA
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cggaggtagg atca
                                                                     14
<210> 70
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ccacccgcct cag
                                                                    13
<210> 71
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cacaggagtg gttgg
                                                                    15
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 <211> 14
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 <223> Description of Artificial Sequence:probe sequence
 <400> 72
cggaccagtg cgtg
                                                                     14
<210> 73
<211> 14
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tcggaccagt gcgt
                                                                     14
<210> 74
<211> 14
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aacggggtac gata
                                                                    14
<210> 75
<211> 18
<212> DNA
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cagttgagat tctaagac
                                                                    18
<210> 76
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aggggcgtct tg
                                                                    12
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<210> 77
<211> 15
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<223> Description of Artificial Sequence:probe sequence
<400> 77
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gtaagtaggc atagc
<210> 78
<211> 13
<212> DNA
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<400> 78
                                                                    13
tgcccagccc cag
<210> 79
<211> 14
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ccaacactcg tgaa
<210> 80
<211> 15
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                                                                    15
gtaagtagac acagc
<210> 81
<211> 12
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<400> 81
                                                                     12
tcggaccagt gc
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<211> 13
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence:probe sequence
<400> 82
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cgatcacgct ggc
<210> 83
<211> 13
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gtcctggggg tgg
<210> 84
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<223> Description of Artificial Sequence:probe sequence
<400> 84
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gtaagtaggt gtgac
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ggttgtacgg gttcacg
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ggaccagtgc gtga
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<210> 87
 <211> 15
 <212> DNA
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gtaagtagac gcagc
                                                                    15
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gtaagtaggc gcagc
                                                                    15
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gtaagtaggc gcggc
                                                                    15
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ggttcccgag cg
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